

Amendments to Specification

PI ase replac paragraph starting on page 3, line 1:

Figure 1A-B depicts the amino acid sequence alignment between the N-Acetylglutamate Kinase from *Synechocystis* sp. (NCBI General Identifier No. 1652434; SEQ ID NO:11), the instant corn contig assembled of clones cr1n.pk0113.g3 and p0111.cipmf81r (SEQ ID NO:2), the instant rice clone rls72.pk0023.c6 (SEQ ID NO:4), the instant soybean clone sdp2c.pk010.h6 (SEQ ID NO:6) and the instant wheat contig assembled of clones wlk1.pk0019.d6 and wl1.pk0009.a7 (SEQ ID NO:8). The top row indicates with asterisks (*) the amino acids conserved among all sequences and with plus signs (+) the amino acids conserved only among the plant sequences. Dashes are used by the program to maximize the alignment of the sequences. The bottom row contains the consensus sequence (SEQ ID NO:12) comprising all the amino acids which are conserved among all plant sequences.

Please remove last two pages of the specification.

The two pages following page 11 of the Sequence Listing are a copy of the International Search Report which, by a clerical error, was attached to the Application as filed.